

Figure 1

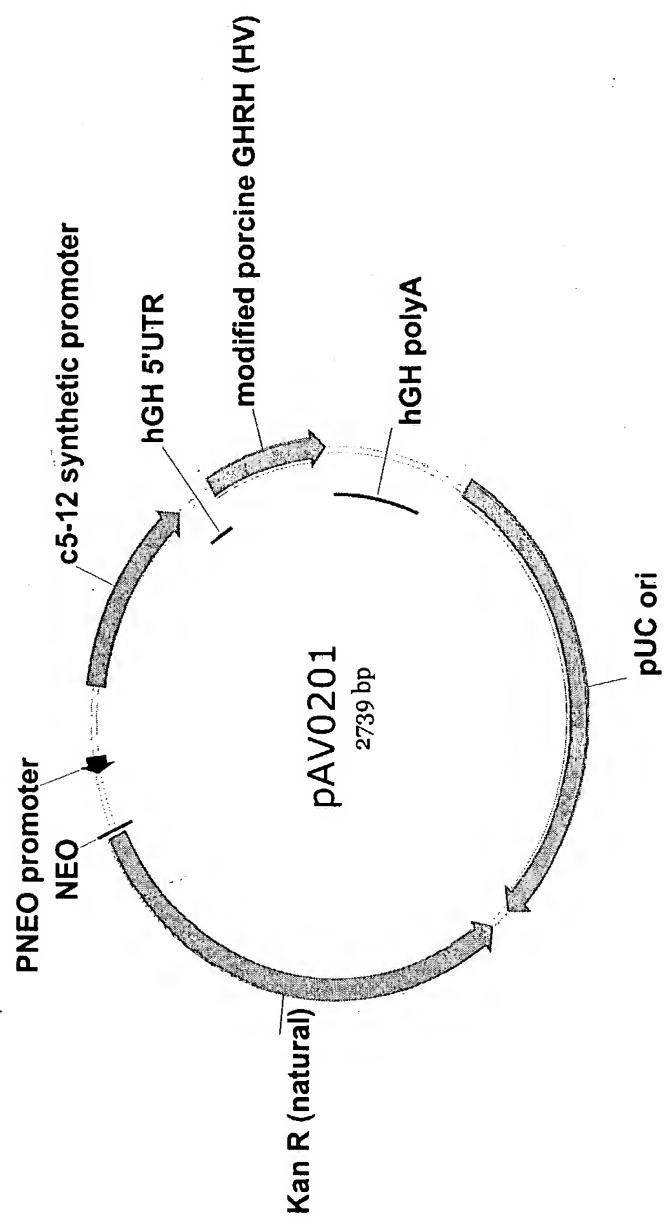


Figure 2

M I E Q D G L H A G S P A A W V E R L F G Y D W A Q Q T I G C S D A .
 1 ATGATTGAAAC AAGATGGATT GCACCGAGGT TCTCGGGCCG CTTGGGTGGA GAGGTACTTC GGCTATGACT GCAATCGGC TGCTCTGATG
 TACTAATG TCTACCTAA CTCGCTTCA AGAGCCGGC GAAACCCRCTT CTCGATTAAG CCGATACCTA CCGCTGTTG CTGTTGCGC AGGAGACTAC
 . A V F R L S A Q G R P V L P V K T D L S G A L N E L Q D E A A R L .
 101 CCGCCGGTGT CCGGCTGTCA GGCAGGGGC GCCCCGGTCTT TTTGGTCAAG ACCGACTGT CCGGTTGCTT GAATGAACTG CAGGAGGAGG CAGGGGGCT
 GGCAGACAA GGCAGACAGT CGGTCTCCG CGGGCAAGA AAGACAGTC TGGCTGAGA GGCAGCGGA CTIACCTAC GTCCTGCTCC GTGGGCCA
 . S W L A T T G V P C A A V L D V V T E A G R D W L L G E V P G Q
 201 ATCGTGGCTG GCCACGAGG GOGTCTTG CGCAGCTTG TCACGCTTG CTCAGCTGAGC GGGAAAGGAC TGGCTGCTAT TGGGGAAAGT GCGGGGCCAG
 TAGAACCGAC CGTGTGTCG CGAACGAAAC GCGTGAACAC GACCTGCAAC AGTGACTCTG CCCTTCCTG ACCGACATA ACCGTTCA CGGGCCGTC
 D L I S S H L A P A E K V S I M A D A M R R L H T L D P A T C P F D .
 301 GATCTCCTGT CATCTCACCT TGCTCTGCG GAGAAAGT CCTCATGCG TGATGGAATG CGGCGCTGC ATAGCTTGA TCCGGTACCC TGCCATTGCG
 CTAGAGGACA GTAGAGTGGA ACAGGAGCGG CTCTTICATA GGTAGTACCG ACTACCTAC GCGCGGAGC TATGGAAC AGGCATGG AGGGTAAGC
 . H Q A K H R I E R A R T R M E A G L V D Q D D L D E E H Q G L A P .
 ACCACAAAGC GAAACATGCG ATCGAGGAG CACGTAATCG GATGGAAGCC GGCTCTGCG ATCGGATGA TCTGGACAA GAGCATAGG GGCTGCGCC
 TGGTGTGTCG CTTGTAGCG TAGCTGGCTC GTGGATGAGC CTACCTGG CGAGAACAGC TAGTCTCTACT AGACCTGCTT CTCGATGTC CCGAACGCGG
 . A E L P A R L K A R M P D G E D L V V T H G D A C L P N I M V E N
 AGCCCAACTG TTGGCCAGGC TOAAGGGCG CATGCCGAC GGGGAGGATC TCGTGTGAC TCATGGCAT GCTGTGCTGC CGAATATCAT GTGAAAAT
 TCGGTTGAC AAGGGTTCG AGTCCCGCG GTACCGGCTG CCCTCTCTAG AGCAGCTG AGTACCGCTA CGGACGAAGC GCTTATAGTA CCACCTTTTA
 G R F S G F I D C G R L G V A D R Y Q D I A L A T R D I A E E L G G .
 401 GGCCTCTT CTGGATTAT CGACTCTGGC CGGCTGGTGTG TGCGGGACCG CTATCAGAC ATAGCTGTTGG CTACCGTGA TATTGCTGAA GAGCTTGGCG
 CCGGGAAA GACCTAAGTA GCTGACACCG GCGAACCCAC ACCGCTGGC GATAGCTCTG TATCCAAAC GATGGCCTT ATAACGACTT CTGAAACCGC
 . E W A D R F L V L Y G I A A P D S Q R I A F Y R L L D E F F *
 501 CGGAATGGGC TGACCGCTTC CTCGTTGCTT ACGGTATGCG CGTCCCGAT TCGCAGGGCA TCGCCTTCTA TCGCCTTCTA GACGAGTCTI TCTGA
 CGCTTACCCG ACTGGCGAAG GAGCACGAAA TGCCATAGCG GCGAGGGCTA AGCGGAAGT AGCGTGGCT AGCGGAAGT AGCGTGGCT AGCGGAAGT
 601

Figure 3

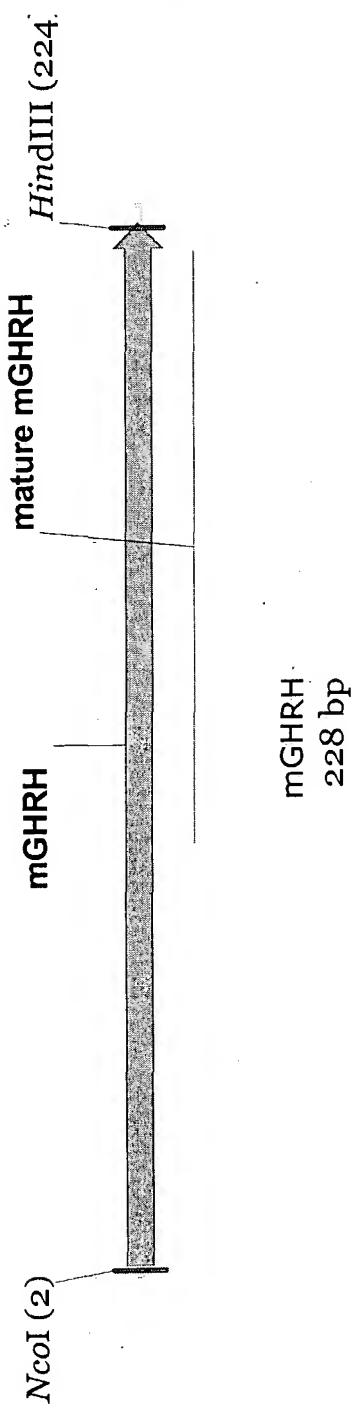


Figure 4

+3 A M V L W V L F V I L I L T S G S H C S L P P S P F R M Q R H V
 1 GCCATGGGC TCTGGGTGCT CTTTGATC CTCATCTCA CCAGGGCAG CCACATGGCAG CTGGCTCCCA GCTCTCCCTT CAGGATGGAG AGGGACCTGG
 CGTACCAAG AGACCCACGA
 +3 D A I F T T N Y R K L L S Q L Y A R K V I Q D I M N K Q G E R I Q E
 101 ACGCCATCTT CACCAAC TAGGAAAC TGCTAGCCA GCTGTACGCC AGGAAGGTGA TCCAGGACAT CATGAAACAG CAGGGCGAGA, GGATCCAGGA
 TGCGGTAGAA GTGGTGGTIG ATGTCCTCG ACGACTCGGT CGACATGCCG TCCTTCCACT AGGTCTGTAA GTACTGTTC GTCAGGTCT CCTAGGTCT
 +3 Q R A R L S & # A C
 201 GCAGAGGGCC AGGCTGAGCT GATAAGCTTG C
 CGTCTCCGG TCCGACTCGA CTATTGAAAC G

Figure 5

GHRH-m	Ori	CCATGGTGTCTGGGTGCTTGTGATCCTCATCCTCACCAGTGGCTCCACTGCTCA	60
GHRH-m	Opt	GCCATGGTGTCTGGGTGCTTGTGATCCTCATCCTCACCAGGGAGCCACTGAGC	
GHRH-m	Ori	CTGCCCCCTCACCTCCCTCAGGATGGCAGGACACGTGGACGCCATCTTCAACCAAC	120
GHRH-m	Opt	CTGCCTCCAGCCCTCCCTCAGGATGGCAGGGCACGTGGACGCCATCTTCAACCAAC	
GHRH-m	Ori	TACAGGAAGCTGCTGAGCCAGCTGAGCCAGGTGATCCAGGACATCATGAACAG	180
GHRH-m	Opt	TACAGGAAGCTGCTGAGCCAGCTGAGCCAGGTGATCCAGGACATCATGAACAG	
GHRH-m	Ori	CAGGGGAGAGAATCCAGGAGGAGGGCCAGGGCTGAGCTGATAAGCTT..	231
GHRH-m	Opt	CAGGGGAGAGGATCCAGGAGGGCCAGGGCTGAGCTGATAAGCTG	

Figure 6

Figure 7

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60

GHRH-M Ori .MVLWVLFVILILTSGSHCSLPPSPPFRMQRHVDIAIFTTNYRKILLSQLYARKVIQDIMNK
GHRH-M opti AMVLWVLFVILILTSGSHCSLPPSPPFRMQRHVDIAIFTTNYRKILLSQLYARKVIQDIMNK

GHRH-M Ori QGERIQEQRARLSA. 75

GHRH-M opti QGERIQEQRARLSAC

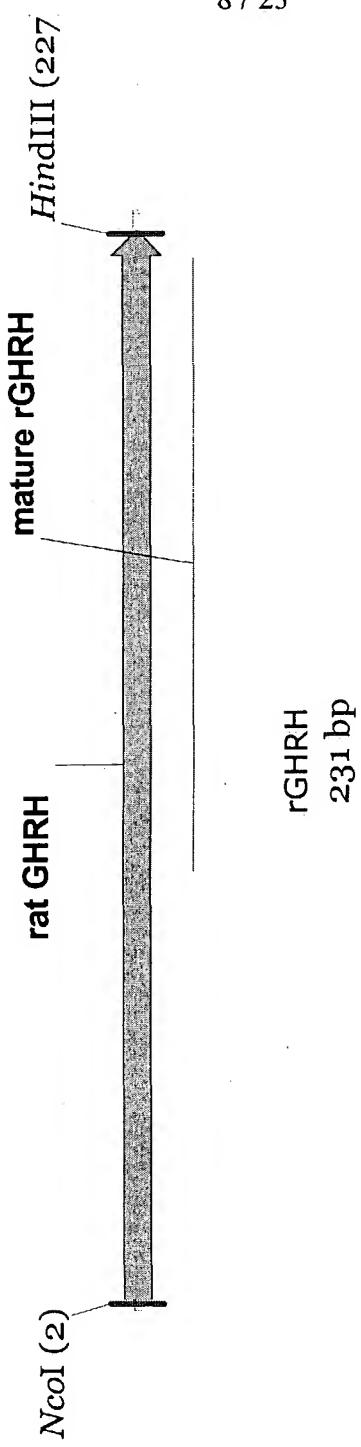


Figure 8

+3 A M A L W V F F V L L T L T S G S H C S L P P S P P F R V R R H A
 1 GGCATGGCC TGTGGGTGTT CTTCGTGTG CTGACCTCTGA CCAGGGAAAG CCACTGGAGC CTGCCTCCCA GCCTCCCTT CAGGGTGCAGC CGGCACGCC
 CGGTACCGGG ACACCCACAA GAAGCACGAC GACTGGGACT GGTGACCTTC GGTGACCTCG GACGGAGGGT CGGGAGGGAA GTCCCACGCG GCGGTGCGGC
 +3 D A I F T S S Y R R I L G Q L Y A R K L L H E I M N R Q Q G E R N Q
 101 ACGCCATCTT CACCAAGCAGC TACAGGAGGA TCCTGGCCA GCTGTACGCT AGGAAGCTC TGCAAGGAGT CATGAACAGG CAGCAGGGG AGAGGAACCA
 TGGGTAGAA GTGGTGTG AGGACCCGGT AGGACCCCT ATGTCCTCTG AGTGTGTGCA TCCCTTCGAGG ACGATGCGA TCCATGCGA GTACTGTCC GTCTCCCTGGT
 +3 E Q R S R F N & # A C
 201 GGAGCAGAGG AGAGGGTCA ACTGATAGC TTGC
 CCTCGTCTCC TCGTCCAGT TGACTATTG AACG

Figure 9

GHRH-R	Ori	GCCATGGCACTCTGGGTGTTCTTGTCTACCCCTACCAAGTGGCTCCACTGCTCA	60
GHRH-R	opti	GCCATGGCCCTGTGGGTGTTCTTGTGCTGTTGACCCCTGACCAGGGAGGCCACTGCGAGC	
GHRH-R	Ori	CTGCCCTCACCTCCCTTCAGGGTGGGGCACGCCATCTTACCCAGCAGC	120
GHRH-R	opti	CTGCCCTCCAGGCCCTCCCTTCAGGGTGGGGCACGCCATCTTACCCAGCAGC	
GHRH-R	Ori	TACAGGAGAATCTGGCAGGTGTAGGCCAGGAAACTGCTGCACGAGATCATGAACAGG	180
GHRH-R	opti	TACAGGAGGATCTGGCAGGTGTAGGCTAGGAAGCTCTGGCAGGAGATCATGAACAGG	
GHRH-R	Ori	CAGCAGGGAGGAAACAGGAGGAGAGGTCCAGGTTCAACTGATAAGCTTGC	234
GHRH-R	opti	CAGCAGGGAGGAAACAGGAGGAGAGGTCCAGGTTCAACTGATAAGCTTGC	

Figure 10

GHRH-R Ori .MALWVFFVLLTSGSHCSLPPSPPFRVRRHADAIFTSSYRRLGQLYARKLILHEIMNR
GHRH-R opti AMALWVFFVLLTSGSHCSLPPSPPFRVRRHADAIFTSSYRRLGQLYARKLILHEIMNR

GHRH-R Ori QQGERNQEQRSRFNA. 76
GHRH-R opti QQGERNQEQRSRFNA.C

Figure 11

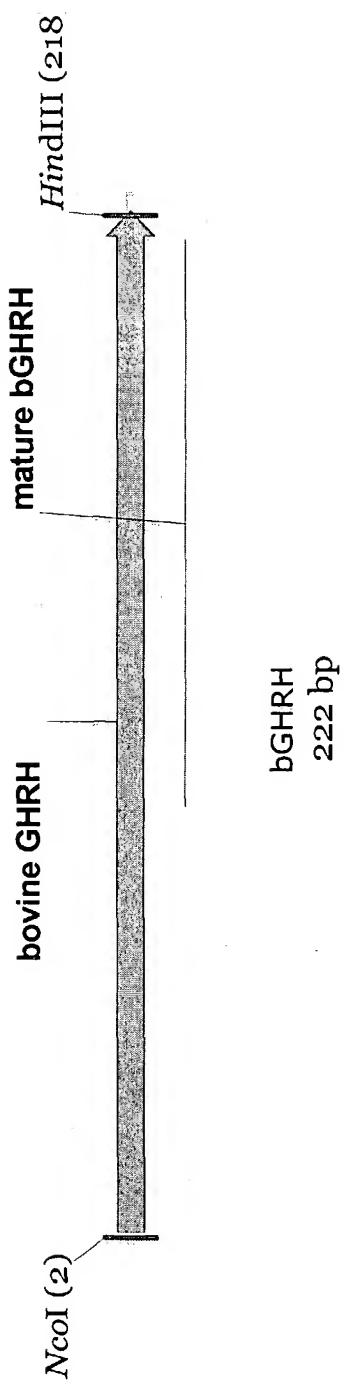


Figure 12

+3 A M V L W V F F L V T L T L S S G S H G S L P S Q P L R I P R Y A
 1 GCCATGGTGC TGTTGGTGT CTCCTGGT ACCCTGACCC TGAGGAGGG CTCCCACGGC TCCCTGGCCT CCCAGCCTCT GGCATCCCT CGCTACGGCG
 CGGTACCAAG ACACCCACAA GAAGGACCAAC TGGGACTGGG ACTCGTGCAC AGGGTGGCG AGGGACGGGA CGGTGGAGA CGGTAGGGA GCGATGGGC
 +3 D A I F T N S Y R K V L G Q L S A R K L L Q D I M N R Q Q G E R N Q
 101 AGGGCATTT CACCAAGAG TACCGCAAGG TGGTGGCCA GCTCAAGGC CGAAAGCTCC TGCAGGGAT CATAAACGG CGCAGGGGG AGCGAACCA
 TGGGTGAA GTGGTAGAA GTGGTAGAA ATGGTGTGCG ATGGGTTC AGAGCCGGT CGAGCTTCC GTACTTGCC GTACTCTGTA AGTGTGGCC TCGGTCCGC TCGGTGGT
 +3 E Q G A & # A C
 201 GGAGCAGGGAA GCCTGATAAG CTTGC
 CCTCGTCCT CGGACTATTG GAACG

Figure 13

GHRH-B	Ori	CCATGGTGCTCTGGTCTTCTCGTGAACCTCACCCCTCAGCAGGGCTCCACGGT	60
GHRH-B	opti	GCCATGGTGCTGTGGTCTTCTCGTGAACCTCACCCCTGAGCAGGGCTCCACGGC	
GHRH-B	Ori	TCCCTGCCTTCCCAGCCTCTCAGGATTCCACGGTACGCCACGCCATCTTCAACAGC	120
GHRH-B	opti	TCCCTGCCTTCCCAGCCTCTGCCATCCCTCGCTACGCCACGCCATCTTCAACAGC	
GHRH-B	Ori	TACCGGAAGGTGCTGGCCAGCTGTCCGCCAGGCTGCTGCAGGACATCATGAACAGG	180
GHRH-B	opti	TACCGCAAGGTGCTGGCCAGCTAGGCCAGGCTAGGCCAGGCTCAAGCTGCAGGACATCATGAACAGG	
GHRH-B	Ori	CAGCAGGGCGAGAGAACCCAGGAGCAGGGGCCGTGATAAGCT.	225
GHRH-B	opti	CAGCAGGGCGAGGCCAACCCAGGAGCAGGGAGCAGCTGATAAGCTTGCG	

Figure 14

60
 GHRH-B Ori .MVLWVFFLVTLLTSSGSHGSLPSQPLRIPRYADALFTNSYRKVLGQLSARKLQLQDIMNR
 GHRH-B opti AMVLWVFFLVTLLTSSGSHGSLPSQPLRIPRYADALFTNSYRKVLGQLSARKLQLQDIMNR
 GHRH-B Ori QQGERNQEQQAA. 73
 GHRH-B opti QQGERNQEQQAAAC

Figure 15

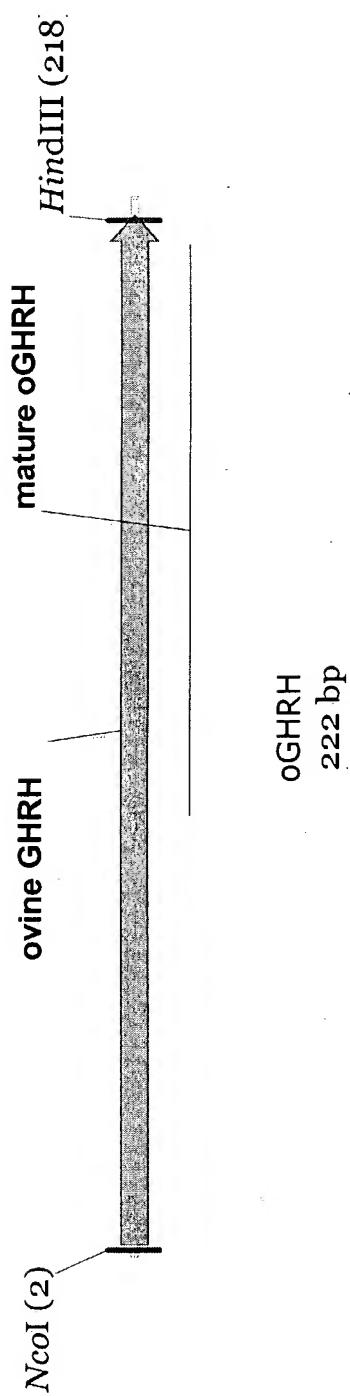


Figure 16

+3 A M V L W V F F L V T L T L S S G S H G S L P S Q P L R I P R Y A
 1 GCCATGGTGC TGTGGTGTG ACCCTGACCC TGAGCAGGG AAGCCACGGC AGCCAGCCCA GCGAGCCCT GAGGATCCCT AGGTACGCC
 CGGTACACG ACACCACAA GAAGGACCA C TGGACTGGG ACTCGTGGC TTGGACTGGG TCGGACGGT CGGTGGGA CTCCTAGGA TCCATGCCC

 +3 D A I F T N S Y R K I L G Q L S A R K L L Q D I M N R Q Q G E R N Q
 101 AGCCCATCTT CACCAACAGC TACAGGAAGA TCCTGGGCCA GCTGAGGGT AGGAAGCTCC TGCAGGACAT CAGAACAGG CAGCAGGGG
 TGCGTAGAA GTGGTGTG ATGTCCTCT AGACCCGGT CGACTCGGA TCCTTCAGG ACGTCTGTA GTACTTGTC GTCGTCGGC TCTCCTGGT

 +3 E Q G A & # A C
 201 GAGGAGGGC GCCTGATAAG CTTGC
 CCTCGTCCG CGGACTATTG GAAAG

Figure 17

GHRH-O	Ori	CCATGGTGTCTGGTGTCTCGTACCCCTCACGCTGGCTCCACGGT	60
GHRH-O	opti	GCCATGGTGTCTGGTGTCTCGTACCCCTGAGGGAAAGCACGGC	
GHRH-O	Ori	TCCCTGCCTTCCCAGCCTCTCAGGATTCCACGGTACGCCGACGCCATCTTCACCAAACAGC	120
GHRH-O	opti	AGCCCTGCCAAGCCAGCC CTGAGGATCCCTAGGTACGCCGACGCCATCTTCACCAAACAGC	
GHRH-O	Ori	TACCGGAAGATACTGGCCAGCTGTCCGCCAGGACTCATGAACAGG	180
GHRH-O	opti	TACAGGAAGATCCTGGCCAGCTGAGGCTAGGACTCATGAACAGG	
GHRH-O	Ori	CAGCAGGGGAGAGAACCGGAGGAGGAGGAGGAGGAGGAGGAGG	225
GHRH-O	opti	CAGCAGGGGGAGAGGAACCAGGAGCAGGGCCTGATAAGCTTGC	

Figure 18

Figure 19

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GHRH-O Ori .MVLWVFFFLVTLTSSGSHGSLPSQPLRIPRYADAIFTNSYRKILGQLSARKLILQDIMNR 60
GHRH-O opti AMVLWVFFFLVTLTSSGSHGSLPSQPLRIPRYADAIFTNSYRKILGQLSARKLILQDIMNR

GHRH-O Ori QQGERNQEQQGAA. 73

GHRH-O opti QQGERNQEQQGAAAC

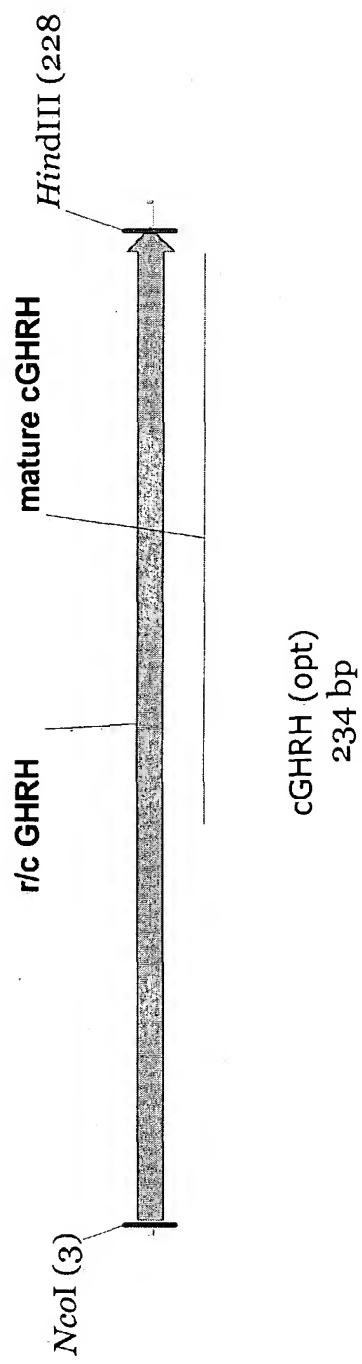


Figure 20

Figure 21

+3 A M A L W V F F V L L T L T S G S H C S L P P S P P F R V R H A
1 GCAATGGCC TGTGGGTCTT CTTTGTGCTG CTGACCCCTGA CCTCCGGAAAG CCACTGCAGC CTGCCACCCA GCCACCCCTT CCGGGTAGG CGCACGCC
CGTACCGGG ACACCCACAA GAAACACGGAC GACTGGGACT GGAGGCCCTTC GGTGACGTG GACGGGGGT CGGGTGGAA GGGGAGTCC GCGGTGGGC
+3 D G I F S K A Y R K L L G Q L S A R N Y L H S L M A K R V G S G L G
101 ACGGCAATCTT CAGCAAGGCC TACCGCAAGC TCCTGGGCCA GCTGAGCGCA CGCAACTACC TGCACAGGCCT GACTCGGT CGGTGATGG AGTGGCAAG CGCTGGCA GCGACTGGG
TGCCGTAGAA GTCGTTCCGG ATGGCGTTCG AGGACCCGGT CGACTCGGT CGGTGATGG ACGTGTCGGA CTACCGTTT GCGCACCCGT CGCTGACCC
+3 D E A E P L S & # A C
201 AGACCGAGGCC GAGCCCCCTGA GCTGATTAAGC TTGC
TCTGCTCGG CTCGGGACT CGACATTCG AACG

Figure 22

Figure 23